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INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(51) International Patent Classification ⁵ : C12Q 1/02	A1	(11) International Publication Number: WO 92/01063 (43) International Publication Date: 23 January 1992 (23.01.92)
(21) International Application Number: PCT/US91/04699 (22) International Filing Date: 2 July 1991 (02.07.91) (30) Priority data: 549,394 6 July 1990 (06.07.90) US (71) Applicant: BIOLOG, INC. [US/US]; 3447 Investment Boulevard, Suite 3, Hayward, CA 94545 (US). (72) Inventor: BOCHNER, Barry ; 2904 Lincoln Avenue, Alameda, CA 94501 (US). (74) Agents: CARROLL, Peter, G. et al.; Limbach, Limbach & Sutton, 2001 Ferry Building, San Francisco, CA 94111 (US).		(81) Designated States: AT (European patent), AU, BE (European patent), CA, CH (European patent), DE (European patent), DK (European patent), ES (European patent), FR (European patent), GB (European patent), GR (European patent), IT (European patent), JP, LU (European patent), NL (European patent), SE (European patent). Published <i>With international search report.</i>
(54) Title: METHODS FOR DETECTION, IDENTIFICATION AND SPECIATION OF LISTERIAS (57) Abstract The present invention relates generally to differential carbon source metabolism in the genus <i>Listeria</i> , metabolic, biochemical, immunological and genetic procedures to measure said differential carbon source metabolism and the use of these produces to detect, isolate and/or distinguish species of the genus <i>Listeria</i> as well as detect, isolate and/or distinguish strains of species of <i>Listeria</i> . The present invention also contemplates test kits and enrichment media to facilitate these procedures.		

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**METHODS FOR DETECTION, IDENTIFICATION
AND SPECIATION OF LISTERIAS**

Field of the Invention

5 The present invention relates to differential carbon
source metabolism in Listeria as a basis for
detection, identification and speciation.

Background of the Invention

10 Bacteria belonging to the genus Listeria are
Gram-positive, non-spore forming, motile rods
characterized in part by their capability for growth
over a wide range of temperatures (4°C to 45°C). H.
Zinsser, Microbiology (14th Edition) (Meredith Corp.
1968) pp. 751-753. The taxonomic relationships
15 between the genus and allied Gram-positive taxa are
not clear. Furthermore, since all Listeria show a
high degree of DNA relatedness and very close
similarity in biochemical, phenotypic and protein
characteristics, there is disagreement in the field
as to relationships between the identified species
20 within the genus. B.J. Wilkinson and D. Jones, J.
Gen. Microbio. 98:399 (1977). J. McLauchlin, J.
Appl. Bacterio. 63:1 (1987).

25 Improved characterization of the genus would
have important medical consequences. Outbreaks of
listeriosis in the United States have been reported
from Listeria-contaminated food. For example, in
1981, there was a major outbreak associated with
cabbage, in 1983 with milk and in 1985 with
contaminated cheese. Listeria infection in humans
30 can cause a variety of symptoms ranging from cold-
like to flu-like but is especially dangerous to a
fetus where it induces a 50% mortality rate.

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With the focus on disease prevention, rapid screening of food samples for Listeria is of foremost concern to the food industry. Under current regulations, the presence of viable cells of any Listeria species in foods is a cause for concern. Thus, there is a need, as a first step, to improve the capability to detect, identify and speciate all species of the genus.

With the need to improve the capability to speciate all species, there is also the need to better understand the pathogenicity, if any, with each particular species. Nearly all of the reported cases of human infections by bacteria belonging to the genus Listeria have been caused by Listeria monocytogenes. See McLauchlin, J. Appl. Bacterio. 63:1 (1987). However, instances have been reported in which Listeria ivanovii, Listeria innocua and Listeria seeligeri have caused disease in humans. It is thus, as a second step, important to determine whether specific strains are associated with disease.

The Food and Drug Administration (FDA) in the United States and comparable agencies in other countries have promulgated standard laboratory methods to detect the presence of Listeria in environmental or food specimens (eg milk). These methods involve culturing an appropriately prepared sample on microbiological media under conditions favorable for growth of these organisms and unfavorable for other bacteria. Detection of bacteria of the Listeria genus is attempted by examining the resulting colonies for morphological and biochemical characteristics, a process that typically is started 48 hours after acquisition of the sample and takes between 9 - 19 days to complete.

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Newer methods of detecting Listeria include a) nucleic acid probes capable of binding to the nucleic acid of particular species, and b) antibodies capable of reacting with antigens specific to particular species. See e.g. A.R. Datta et al., Appl. Environ. Microbio. 54:2933 (1988) (probes to a fragment of a presumptive hemolysin gene of L. monocytogenes); J. Klinger, J. Assoc. Off. Anal. Chem 71:669 (1988) (nucleic acid hybridization assay for Listeria in foods); European Patent Application No. 0355147 (corresponding to U.S.S.N. 227,402 and 143,490) (a probe directed to the hemolysin gene in L. monocytogenes); European Patent Application No. 0314294 (corresponding to U.S.S.N. 965,510) (probes to rRNA of L. monocytogenes); G.R. Siragusa and M.G. Johnson, Appl. Environ. Microbio. 56:1897 (1990) (monoclonal antibodies to an antigen shared with three Listeria species); J.A. Mattingly, J. Assoc. Off. Anal. Chem. 71:679 (1988) (antibody-based assay for detection of L. monocytogenes); European Patent Application No. 0303309 (corresponding to U.S.S.N. 836,619) (antibodies to heat-treated extracts of Listeria).

The present invention now provides a method capable of 1) detecting bacteria of the genus Listeria and 2) distinguishing between species of the genus Listeria. The method of the present invention has the advantage over previous methods in that it is easier, simpler and less expensive. It is also faster than the current biochemical characterization procedures which involve slow fermentation reactions detected with pH indicators.

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Summary of the Invention

In one embodiment, the present invention provides a method for detecting a species of the genus Listeria comprising exposing a sample suspected of containing microorganisms belonging to the genus Listeria to a metabolically effective amount of one or more carbon sources capable of being metabolized by a subset of species of Listeria, for a time and under conditions sufficient for the subset to metabolize the carbon source, and detecting any metabolism of the carbon source by assaying for respiration by the microorganisms, thereby determining the presence of the subset of species.

In another embodiment, the method further comprises, prior to the exposing step, subjecting the sample to an enrichment treatment which will enhance the population of the genus Listeria in the sample.

In another embodiment, the carbon sources are selected from the group consisting of mannitol, D-gluconic acid, L-malic acid, 5'AMP, β -methyl glucoside, glycerol, D-xylose, L-rhamnose, xylitol, sucrose, α -D-lactose, L-lactic acid, and hexose-phosphate.

In another embodiment, the carbon sources comprise a panel of carbon sources comprised of L-lactic acid and xylitol.

In another embodiment, the carbon sources comprise a panel of carbon sources comprised of hexose-phosphate and xylitol.

The present invention also contemplates a test kit for the detection of a species of the genus Listeria or a strain of a species of the genus Listeria comprising at least one compartment containing at least one carbon source capable of being metabolized by a subset of species of the genus Listeria, wherein the carbon source is selected from

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the group consisting of mannitol, D-gluconic acid, L-malic acid, 5'AMP, β -methyl glucoside, glycerol, D-xylose, L-rhamnose, xylitol, sucrose, α -D-lactose, L-lactic acid, and hexose-phosphate.

5 The present invention also contemplates a test kit for the detection of a species of the genus Listeria or a strain of a species of the genus Listeria comprising a plurality of compartments containing a plurality of carbon sources capable of
10 being metabolized by a subset of species of the genus Listeria, wherein the carbon sources comprise a panel of carbon sources, each carbon source of said panel in separate compartments, comprising L-lactic acid and xylitol.

15 The present invention also contemplates a test kit for the detection of a species of the genus Listeria or a strain of a species of the genus Listeria comprising a plurality of compartments containing a plurality of carbon sources capable of
20 being metabolized by a subset of species of the genus Listeria, wherein the carbon sources comprise a panel of carbon sources, each carbon source of said panel in separate compartments, comprising hexose-phosphate and xylitol.

25 The present invention further provides a method for detecting a species of the genus Listeria comprising exposing a sample suspected of containing microorganisms belonging to the genus Listeria to a metabolically effective amount of one or more carbon
30 sources capable of being metabolized by L. monocytogenes and L. ivanovii but not for metabolism by other species of Listeria, for a time and under conditions sufficient for metabolism by L. monocytogenes and L. ivanovii, and detecting any
35 metabolism of said carbon source, thereby determining the presence of L. monocytogenes and/or L. ivanovii.

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In one embodiment of this method, the method further comprises, prior to the exposing step, subjecting the sample to an enrichment treatment which will enhance the population of the genus Listeria in the sample.

In one embodiment of this method, the carbon sources are selected from the group consisting of L-lactic acid and hexose-phosphate.

The present invention also contemplates a method for detecting a species of the genus Listeria comprising exposing a sample suspected of containing microorganisms belonging to the genus Listeria to a metabolically effective amount of one or more carbon sources capable of being metabolized by a subset of species of the genus Listeria, but not metabolized by L. monocytogenes or L. ivanovii, for a time and under conditions sufficient for metabolism by the subset of species, and detecting any metabolism of the carbon source, thereby determining the presence of a species other than L. monocytogenes and L. ivanovii.

In one embodiment of this method, the method further comprises, prior to the exposing step, subjecting the sample to an enrichment treatment which will enhance the population of the genus Listeria in the sample.

In one embodiment of this method, the carbon sources are selected from the group consisting of mannitol, D-gluconic acid, and L-malic acid.

In another embodiment of this method, the carbon sources comprise a panel of carbon sources comprised of mannitol, D-gluconic acid, and L-malic acid.

The present invention also contemplates an enrichment medium for species of the genus Listeria comprising, in liquid or semi-solid form, a basal medium having all the constituents selective for growth of the genus Listeria, and a growth effective

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amount of one or more carbon sources capable of being utilized by a subset of species of the genus Listeria, wherein the carbon source is selected from the group consisting of mannitol, D-gluconic acid, L-malic acid, 5'AMP, β -methyl glucoside, glycerol, D-xylose, L-rhamnose, xylitol, sucrose, α -D-lactose, L-lactic acid, and hexose-phosphate.

In the sense that, heretofore, no known method of detecting the genus Listeria allowed for complete speciation, one aspect of the present invention is an improvement. In a method for detecting microorganisms of the genus Listeria wherein a sample suspected of containing microorganisms belonging to the genus Listeria is subjected to an enrichment treatment capable of enhancing the population of the genus Listeria in the sample, the improvement comprises, thereafter exposing the sample to a metabolically effective amount of one or more carbon sources capable of being metabolized by a subset of species of Listeria, for a time and under conditions sufficient for the subset to metabolize the carbon source, and detecting any metabolism of the carbon source by assaying for respiration by the microorganisms, thereby determining the presence of the subset of species.

Detailed Description of the Invention

The present invention is predicated in part on the surprising discovery that microorganisms of a species of the genus Listeria can be distinguished from microorganisms of another species of the genus Listeria as well as from non-Listeria microorganisms by differential metabolism of a variety of novel carbon sources. The present invention is also based on the surprising discovery that specific strains of species of the genus Listeria can be distinguished

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from other strains of the same species by differential metabolism of a variety of carbon sources.

5 The present invention contemplates that the differential metabolism can be observed at the biochemical level by measuring parameters such as respiration (e.g. measured colorimetrically) and microbial growth (e.g. measured turbidimetrically). The present invention also contemplates that this
10 differential metabolism can also be observed a) using an indicator strain of Listeria that is competent for transformation, b) using nucleic acid probes, or c) employing specific antibodies.

15 Unless otherwise specified, the generic term "Listeria" is used herein to refer to all species of the genus including, but not limited to, L. monocytogenes, L. ivanovii, L. innocua, L. welshimeri, L. seeligeri, L. grayi, and L. murrayi.

20 With respect to the phrase "microorganisms suspected of being of the genus Listeria" it is not meant that the microorganisms must be homogeneous. It may comprise a heterogeneous mixture of Listeria species or strains of one or more species of Listeria and may contain microorganisms not belonging to the
25 genus Listeria. On the other hand, it may comprise homogenous and/or substantially pure cultures of one or more species or strains of Listeria.

30 The term "sample" in the present specification and claims is used in its broadest sense. On the one hand it is meant to include a specimen or culture. On the other hand, it is meant to include both biological and environmental samples.

35 Biological samples may be animal, including human, fluid or tissue, food products and its ingredients such as dairy, vegetable and meat produce, and waste. Environmental samples include

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environmental material such as surface matter, soil, water and industrial samples.

Whether a biological or environmental sample, a sample suspected of containing Listeria may or may not first be subjected to an enrichment means to create a culture of microorganisms suspected of being of the genus Listeria. By enrichment means or enrichment treatment the present invention contemplates i) conventional techniques for isolating Listeria away from other microorganisms and ii) novel techniques for isolating Listeria away from other microorganisms involving the use of one or more of the carbon sources of the present invention. Importantly, it is not intended that the present invention be limited only to one enrichment step or type of enrichment means. For example, it is within the scope of the present invention to, following subjecting a sample to a conventional enrichment means, subjecting the resulting preparation to further purification such that pure or substantially pure cultures of the Listeria species or strain of species are produced.

With respect to a "culture" it is not meant that the enrichment means must only be in a liquid phase. The present invention contemplates a variety of types of enrichment means (whether in or on a liquid or semi-solid medium).

The carbon compounds of choice in accordance with the present invention are those which some species of the genus Listeria can utilize as a carbon source for growth and/or respiration but which are unable or substantially unable to be metabolized by other species of the same genus.

In one embodiment, the present invention involves using L-lactic acid and/or a hexose phosphate (e.g. glucose-phosphate, mannose-phosphate

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and fructose-phosphate) as the carbon sources of choice. Where a hexose phosphate is used, glucose-1-phosphate is a preferred sugar phosphate (primarily because of its stability, purity and commercial availability). Such carbon sources have heretofore never been reported useful for Listeria detection or speciation.

In another embodiment, the present invention also describes the use of xylitol as a carbon source. Xylitol (to be distinguished from D-xylose) is a carbon source heretofore never reported useful for Listeria detection or speciation.

In another embodiment, the present invention also describes the use of adenosine 5'-monophosphate (5'AMP) as a carbon source. 5'AMP is a carbon source heretofore never reported useful for Listeria detection or speciation.

In another embodiment, the present invention also describes the use of glycerol, β -methyl glucoside, D-gluconic acid and/or L-malic acid as carbon sources. Heretofore, it was reported that all of the species could utilize glycerol and α -methyl glucoside, that none of the Listeria species could utilize gluconic acid, and that malic acid was primarily utilized by L. monocytogenes. J. General Microbio. 98:399 (1977). Our findings are not in agreement with many aspects of this report. In the case of malic acid, the report indicates that a mixture of the D- and L-isomers was used. The usefulness of β -methyl glucoside for speciating Listerias has not heretofore been reported.

Finally, the present invention describes the use of the above-named carbon sources together with one or more carbon sources heretofore tested in the detection and/or identification of Listeria (e.g., mannitol, D-xylose, sucrose, α -D-lactose and L-

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5 rhamnose). This is done, however, with the understanding that it is not intended that the present invention be limited to the use of known useful Listeria carbon sources in conjunction with the novel sources described earlier. Furthermore, the present invention extends to chemical, biochemical, biological and/or functional equivalents or homologues of the aforementioned carbon sources including any structural modifications thereto
10 provided that said equivalents, homologues or modified carbon sources are utilized by some species of the genus Listeria. The carbon sources of the present invention may also be added as its corresponding salt. Accordingly, reference to L-lactic acid, for example, includes its corresponding
15 salt forms.

The carbon sources may be used singularly or in combination with other carbon sources. However, in some circumstances, two or more carbon sources may
20 not be appropriate in combination due to catabolite repression or utilization by other non-Listeria organisms.

Antibiotics or other selective agents (e.g. lithium salts) may be used in conjunction with the present invention to inhibit non-Listeria organisms and/or to promote growth of Listeria bacteria.
25

A preferred embodiment of the present invention contemplates a method for detecting a species or the genus Listeria in a biological sample comprising, subjecting said sample to an enrichment means so as to create a culture of microorganisms suspected of being of the genus Listeria, exposing said culture separately or together to amounts of carbon sources effective for metabolism, wherein the carbon sources
30 are selected from the group consisting of L-lactic acid, glucose-1-phosphate, xylitol, D-gluconic acid,
35

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L-malic acid, 5'AMP, glycerol, and β -methyl glucoside, and wherein the exposure is for a time and under conditions sufficient for metabolism, and then detecting metabolism.

5 Another preferred embodiment of the present invention contemplates a method for detecting a species of the genus Listeria in a biological sample comprising, subjecting said sample to an enrichment means so as to create a culture of microorganisms
10 suspected of being of the genus Listeria, exposing said culture to a panel of carbon sources consisting of mannitol, L-lactic acid, glucose-1-phosphate, xylitol, 5'AMP, β -methyl glucoside, L-rhamnose and D-xylose, wherein the carbon sources are in amounts
15 effective for metabolism and wherein said exposure is for a time and under conditions sufficient for metabolism, and detecting metabolism.

 In the practice of the present invention it has been found that L-lactic acid and glucose-1-phosphate
20 as carbon sources are useful to differentiate L. monocytogenes and L. ivanovii from L. innocua, L. welshimeri, L. seeligeri, L. grayi, and L. murrayi; xylitol as a carbon source is useful to differentiate L. monocytogenes, L. innocua, L. welshimeri and L. seeligeri,
25 from L. ivanovii, L. grayi, and L. murrayi; 5'AMP, D-xylose, mannitol, D-gluconic acid and L-malic acid as carbon sources are useful to differentiate L. grayi and L. murrayi from L. monocytogenes, L. ivanovii, L. innocua, L. welshimeri
30 and L. seeligeri; L-rhamnose as a carbon source is useful to distinguish L. monocytogenes, L. innocua and L. welshimeri, from other species of Listeria. Lastly, glycerol and β -methyl glucoside as carbon sources are useful to distinguish L. ivanovii and L. grayi
35 from other species of Listeria since they give

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a negative or a very weak positive reaction with these. (See Table 1).

5 In accordance with one embodiment of the present invention, the carbon sources are added to a reaction vessel containing basal medium and other constituents as required for growth. "Basal medium" as used herein refers to a medium suitable for growth of Listeria but which does not contain sufficient amounts of a carbon source for metabolism. More
10 conveniently, the carbon sources are added separately or in combination into a series of reaction vessels such that a range of carbon sources and/or various combinations of carbon sources can be tested at one time. Alternatively, the reaction vessel(s) may
15 contain the required carbon source to which basal medium is then added or the carbon source may be added to the basal medium already in the reaction vessel. The order of addition of the components in accordance with the present invention is not
20 critical. In addition to the carbon source and basal medium and other components necessary for growth, each reaction vessel may also contain one or more indicator molecules such as, but not limited to, a redox indicator (e.g. tetrazolium), a pH indicator,
25 and various dyes and the like. These approaches are fully described in U.S. Patents Nos. 4,129,483 and 4,235,964 to Barry R. Bochner, hereby incorporated by reference. A generalized indicator useful for practice of the present invention is also described
30 by Bochner and Savageau. See B. Bochner and M. Savageau, Applied and Environmental Microbiology, 33:434 (1977).

35 Analysis of differential carbon source metabolism as contemplated herein is conveniently conducted using a test kit. In a preferred embodiment of the present invention, the test kit

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comprises a microtiter plate containing an appropriate basal medium. The required carbon source is then added to each compartment at the appropriate concentration. Each compartment may, along with the basal medium, also contain an appropriate indicator or the compartment may be so constructed that growth is measurable turbidmetrically. Alternatively, the indicator may be added separately. In any event, the preferred use of the basal medium, carbon source, indicator and any other chemical (or mixture of chemicals) is in a dry form in the microplate.

It is preferred in this case, that the microorganisms suspected to be Listeria are prepared as required in saline or other suitable diluent and then each compartment of the plate inoculated.

Plates useful in the practice of the present invention as a kit are commercially available from Biolog, Inc., Hayward, California. For example, the Biolog MT MicroPlate™ is particularly useful. The MT plate is a 96-well microplate designed to test the ability of an inoculated microorganism suspension to utilize (oxidize) a panel of different carbon sources. Each well of the panel contains a tetrazolium redox dye and a buffered nutrient medium that has been developed and optimized for a wide variety of bacteria. If used in the practice of the present invention, the carbon sources of the present invention may be added to the MT plates either before or after inoculating with a microorganism suspension. For example, with a Biolog MT MicroPlate™, about 0.6 mg of carbon source (e.g. 15 ul of a 4% stock solution) may be added to each well giving a resulting concentration of 0.4% (w/v). After addition the carbon source may be dried. The Biolog's MT MicroPlate™ may be arranged so that there is one or more negative control wells with no carbon

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source and/or one or more positive control wells containing a carbon source (e.g., salicin) or a mixture of carbon sources such that all Listeria strains will give a positive reaction. The remaining wells of a set contain individual carbon sources or a combination of carbon sources that may or may not be metabolized. After inoculation, the reaction vessel, compartment or indicator plate is then incubated at an appropriate temperature or range of temperatures, preferably between 25 and 40°C and most preferably between 30 and 37°C for 1 to 24 hours. Many of the carbon source reactions are apparent even by 1 hour.

As noted earlier, a sample suspected of containing Listeria may be first subjected to an enrichment means and this enrichment means may involve i) conventional techniques for isolating Listeria away from other microorganisms or ii) novel techniques for isolating Listeria away from other microorganisms involving the use of one or more of the carbon sources of the present invention.

Where conventional techniques are used in the enrichment step, a sample suspected to contain one or more species of Listeria may be inoculated onto a semi-solid medium (e.g. an agar plate) as an enrichment means. The inoculated semi-solid medium is then optionally exposed to conditions which allow for growth of the Listeria. Conventional enrichment means may be selected from any available to those skilled in the art, including but not limited to, the following: original McBride Listeria agar formulations, modified McBride Listeria agar formulation, LiCl-phenylethanol-moxalactam (LPM) agar, acriflavine-ceftazidime agar, Rodriguez isolation agar (RISA), modified Vogel-Johnson (MVJ) agar and cyclohexanedione-nalidixic acid-phenylethanol agar and their liquid forms. Most

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preferably, the enrichment means is LPM or MVJ. A
general discussion of other suitable media for
Listeria can be found in Loessner et al., Appl.
Environ. Microbiol. 54: 3003-3007, 1988; Buchanan et
5 al., Appl. Environ. Microbiol. 55: 599-603, 1989;
Lachica, Appl Environ. Microbiol. 56: 167-169, 1990.

Where novel techniques of the present invention
are used in the enrichment step, a sample suspected
to contain one or more species of Listeria may be
10 inoculated onto a semi-solid medium (e.g. an agar
plate). The inoculated semi-solid medium is then
optionally exposed to conditions which allow for
growth of the Listeria. In any event, microorganisms
present on the plate may then be replica-plated or
15 transferred by other means to liquid or semi-solid
growth media containing a carbon source of the
present invention which may be utilized by one or
more species of Listeria but not by other species of
the same genus.

Alternatively, using novel enrichment techniques
of the present invention, the sample may be
inoculated directly into or onto a liquid or semi-
solid growth medium containing, as sole carbon
source, a carbon source of the present invention
25 capable of being utilized by one or more species of
Listeria but not by other species of the same genus.

In one embodiment, utilization of a carbon
source is measured by the formation of one or more
colonies of the pathogenic strain. The amount of
30 carbon source added to the reaction vessel will be
that required for growth of the Listeria bacterium.
By "growth" it is meant that the bacterium has
undergone at least one doubling, the only requirement
being that sufficient doubling must take place in
35 order for the particular indicator to detect growth,

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or for there to be sufficient growth to be measured turbidimetrically. When a semi-solid growth medium is used, sufficient growth is required to observe one or more colonies. Typically, the final concentration of carbon source is from about 0.05 to about 5.0% (w/v) and preferably from about 0.20 to about 1.0% (w/v). Most preferably, the concentration of carbon source is approximately 0.4% (w/v). However, the concentration may be varied depending on the organism to be detected, availability of carbon source, ease of solubility and indicator selected.

According to this embodiment of the present invention, there is provided an enrichment or selection growth medium for one or more species of Listeria comprising in liquid or semi-solid form, a basal medium having all the constituents required for growth of Listeria except for a carbon source and a growth effective amount of one or more carbon sources capable of being utilized by one or more species of Listeria but not by other species of the same genus.

The present invention also contemplates a novel enrichment means wherein the aforementioned conventional media are used, but wherein the principle carbon source is replaced by one or more of the carbon sources of the present invention.

Other Detection Methods

As noted earlier, the present invention contemplates that the differential metabolism can be observed at the metabolic or biochemical level by measuring parameters such as respiration (e.g. measured colorimetrically) and microbial growth (e.g. measured turbidimetrically). On the other hand, the present invention contemplates that this capability for differential metabolism can also be observed by other detection methods including a) using an

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indicator strain of Listeria in a genetic repair assay, b) using nucleic acid probes in a hybridization assay, or c) employing specific antibodies in an immunological assay or stain.

5 a) Genetic Repair Assay

Another assay contemplated by the present invention allows the rapid screening of biological samples for the presence or absence of pathogenic strains of Listeria. The assay provides a simple
10 yes/no test whether or not a biological sample, such as food, contains a pathogenic strain of Listeria.

In one embodiment, an indicator strain of Listeria is rendered competent for transformation by naked DNA and also carries a mutation affecting
15 expression of one or more enzymes or other proteins involved in the uptake or metabolism of a carbon source by pathogenic strains of Listeria and which carbon sources are not metabolized by non-pathogenic strains of Listeria (examples of such carbon sources
20 may be L-lactic acid or hexose-phosphate; see Table 1 and accompanying discussion, below). The indicator strain may also express a selective marker such as but not limited to resistance to an antibiotic or other antimicrobial compound including a heavy metal.
25 The biological sample is then subjected to genetic extraction meaning that the DNA of any microorganism therein is released by such means as enzymes, organic extraction, detergents or sonic disruption. Further purification and partial digestion of the genetic
30 extract may then be required depending on the biological sample, time and degree of sensitivity desired.

The genetic extract is then mixed with an excess of the competent indicator strain for a time and
35 under conditions sufficient to allow transformation

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of the indicator strain with any Listeria DNA. The mixture is then added dropwise or by other means to the semi-solid or liquid medium. The mixture is introduced to the medium containing, as sole carbon source, the carbon source on which the mutation it carries prevents it from growing. The medium is then incubated and examined for growth. If growth does occur, the indicator strain must have been transformed with DNA from a pathogenic strain and the mutation preventing the indicator strain from growing on the sole carbon source in the medium repaired. Appropriate controls are also run to verify that the genetic lesion did not revert and that the sample extract containing DNA was not contaminated with viable microorganisms.

Many variations to the above assay may be made without departing from the scope of the present invention. For example, rather than the DNA entering the indicator strain by transformation, it might enter by transduction following an in vitro phage packaging reaction, or by conjugation, possibly with a non-Listeria competent strain that would take up naked DNA and transfer it by conjugation into the Listeria indicator strain. In the latter case, the growth medium may additionally contain an agent to prevent growth of the non-Listeria bacterium. Transformation techniques are disclosed, for example, in U.S. Patents Nos. 3,930,956 and 4,038,143 to E. Juni, hereby incorporated by reference. Other approaches are disclosed in A. Janick et al., J. Clin. Microbio. 4:71 (1978). Transformation techniques for Listeria are disclosed in Lucas and Levin, Letters in Applied Microbiology 9:215 (1989).

Accordingly, this aspect of the present invention contemplates a means for detecting a potentially pathogenic strain of a species of the

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genus Listeria in a biological or environmental sample comprising 1) providing a competent strain of Listeria carrying a mutation in or near a gene encoding an enzyme or other protein or parts thereof involved in or associated with the uptake or metabolism of a carbon source used by a potentially pathogenic strain of Listeria, or in or near a promoter region, regulatory gene or other control sequence or parts thereof for expression of said enzyme or other protein, said mutation resulting in said enzyme or other protein not being produced or being produced in inactive form, thereby said competent strain of Listeria being unable to grow on said carbon source, 2) contacting said competent strain of Listeria carrying said mutation with a genetic extract of said biological or environmental sample for a time and under conditions sufficient for said competent strain of Listeria to be transformed with DNA of a potentially pathogenic strain of Listeria, if present, in said genetic extract of the sample, wherein said DNA is capable of repairing the mutation in said competent strain of Listeria and 3) detecting growth of said transformed Listeria on the carbon source for which previously the strain could not grow.

Another aspect of the present invention provides a means for detecting a potentially pathogenic strain of a species of the genus Listeria in a biological or environmental sample comprising 1) providing an indicator strain of Listeria carrying a mutation in or near a gene encoding an enzyme or other protein or parts thereof involved in or associated with the metabolism of a carbon source used by a potentially pathogenic strain of Listeria, or in or near a promoter region, regulatory gene or other control sequence or parts thereof for expression of said

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enzyme or other protein, said mutation resulting in said enzyme or other protein not being produced or being produced in inactive form, thereby said indicator strain of Listeria being unable to grow on said carbon source, 2) contacting said indicator strain of Listeria carrying said mutation with a conjugative donor strain or a phage for a time and under conditions sufficient for said indicator strain of Listeria to be conjugated or transduced with DNA from a potentially pathogenic strain of Listeria, if present, in said sample, wherein DNA transferred is capable of repairing the mutation in said indicator strain of Listeria, and 3) detecting growth of said repaired Listeria on the carbon source for which previously the strain could not grow.

b) Hybridization Assay

Another embodiment of the present invention relates to a nucleic acid molecule capable of hybridizing to genetic material contained in a genetic extract of one or more species of the genus Listeria but which does not hybridize to genetic material in other species of the same genus. Ideally, the nucleic acid molecule will hybridize under predetermined stringency levels to genetic material, at least part of which, encodes an enzyme or protein or portions thereof or defines a promoter region, regulatory gene or other control sequence or parts thereof for expressing enzymes or other proteins associated with the metabolism by particular species of Listeria of the aforementioned carbon source compounds, said genetic material substantially absent in other species of the same genus.

In this regard, the genetic material may be DNA or RNA. When the genetic material is mRNA, the bacterial cells may first need to be inoculated in

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the presence of the subject carbon source for a time and under conditions sufficient to permit transcription of the DNA into mRNA. A genetic extract comprises the genetic material (i.e. DNA or RNA) or various parts thereof from a particular cell.

"Hybridization" is used herein in its broadest sense and refers to the formation of duplexes between completely or partially complementary nucleotide sequences by Watson-Crick base-pairing. The stringency conditions used may vary depending on the base composition of the probe-target duplex as well as by the level and geometry of mispairing between the two nucleic acids.

The genetic detection of a species of Listeria may be accomplished in a number of ways. In one preferred embodiment, the species or strain to be identified is grown on semi-solid media into one or more colonies. The colonies are transferred onto nitrocellulose paper or other suitable material and subjected to dot blot analysis or similar hybridization procedure. Techniques useful for this type of genetic screening can be found in Molecular Cloning: A Laboratory Manual, Sambrook et al eds, Cold Spring Harbor Laboratory, Cold Spring Harbor, second edition, 1989.

The nucleic acid probes may be prepared by any number of means. For example, an enzyme or protein involved in or associated with the metabolism of a carbon source by one or more species of Listeria and which is absent from other species of the same genus is first purified and a contiguous series of amino acids in said enzyme or other protein determined. For example, the N-terminal amino acid sequence is ascertained and a probe comprising nucleotide sequence is prepared based on the amino sequence. Generally a range of probes is produced. Using

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Southern or Northern blots, a probe capable of hybridizing to genetic material in one or more species of Listeria but not genetic material from other species is selected.

5 Conveniently, a biological sample may be screened for the presence of a species of Listeria by immobilizing bacteria contained in said biological sample onto a semi-solid support and then subjecting said immobilized bacteria to detection means by using
10 a labelled nucleotide probe. The label may be a radioactive isotope, or biotinylated molecule or other detectable marker.

 This aspect of the present invention is conveniently provided in kit form comprising a
15 compartment or a multiplicity of compartments adapted to receive a biological sample suspected to contain a species of Listeria, a second compartment or multiplicity of compartments adapted to contain one or more nucleic acid molecules capable of hybridizing
20 at pre-determined stringency conditions to genetic material encoding one or more enzymes or other protein or portions thereof or defining a promoter region, regulatory gene or other control sequence or parts thereof for expressing enzymes or other
25 proteins involved in or associated with the metabolism of one or more carbon sources by one or more species of Listeria and which genetic material is absent or substantially absent from another species of the same genus, said nucleic acid
30 molecules labelled with a reporter molecule.

 In further accordance with this aspect of the present invention, the sensitivity and/or ease of performing this assay may be enhanced by using the polymerase chain reaction (PCR) as described by K.B. Mullis et al. in U.S. Patents Nos. 4,683,195 and
35 4,683,202, hereby incorporated by reference. To this

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end, the present invention extends to nucleotide probes or primers useful in the PCR to detect the aforementioned DNA or RNA. Accordingly, the kit contemplated herein may contain additional compartments adapted to contain the components for PCR including the nucleotide primers.

c) Immunological Assay

A further aspect of the present invention relates to the immunological detection of a species of Listeria using antibodies directed to an enzyme or other protein present in said species but absent in other species of the same genus and which are involved in or associated with the utilization of one or more carbon sources, said enzyme or protein absent in said other species of the same genus. The enzyme or protein is purified from the species by conventional techniques and used to prepare specific antibodies. Such antibodies are useful in developing detection assays (immunoassays) for the enzyme or protein. The presence of the enzyme or protein in a biological sample or an extract thereof is indicative of the presence of a species of Listeria.

In the following methods, where specific enzymes or proteins are not secreted from the bacterium to be detected, the cells are disrupted by treatments which include sonic disruption, osmotic change or use of agents such organic solvents, detergents, enzymes and the like.

Furthermore, immunological equivalents of the enzymes or proteins may be used to facilitate the production of antibodies. Additionally, a mixture of enzymes or proteins may be used to facilitate the production of antibodies. A mixture of enzymes or proteins may also be used to develop antibodies.

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The antibodies may be monoclonal or polyclonal. It is within the scope of this invention to include any second antibodies (monoclonal or polyclonal) directed to the first antibodies discussed above.

5 Both the first and second antibodies may be used in the detection assays or a first antibody may be used with a commercially available anti-immunoglobulin antibody. An antibody as contemplated herein

10 includes any antibody specific to any region of an enzyme or other protein exclusively involved in the metabolism of a carbon source by one or more species of Listeria and which is absent from another species of the same genus.

Both polyclonal and monoclonal antibodies are

15 obtainable by immunization with the enzyme or protein and either type is utilizable for immunoassays. The methods of obtaining both types of sera are well known in the art. Polyclonal sera are less preferred but are relatively easily prepared by injection of a

20 suitable laboratory animal with an effective amount of the purified enzyme or protein, or antigenic parts thereof, collecting serum from the animal, and isolating specific sera by any of the known immunoadsorbent techniques. Although antibodies

25 produced by this method are utilizable in virtually any type of immunoassay, they are generally less favored because of the potential heterogeneity of the product.

The use of monoclonal antibodies in an

30 immunoassay is particularly preferred because of the ability to produce them in large quantities and the homogeneity of the product. The preparation of hybridoma cell lines for monoclonal antibody production derived by fusing an immortal cell line

35 and lymphocytes sensitized against the immunogenic preparation can be done by techniques which are well

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known to those who are skilled in the art. (See, for example Douillard and Hoffman, Basic Facts about Hybridomas, in Compendium of Immunology Vol II, ed. by Schwartz, 1981; Kohler and Milstein, Nature 256: 495-499, 1975; European Journal of Immunology 6: 511-519, 1976).

Unlike preparation of polyclonal sera, the choice of animal is dependent on the availability of appropriate immortal lines capable of fusing with lymphocytes. Mouse and rat have been the animals of choice in hybridoma technology and are preferably used. Humans can also be utilized as sources for sensitized lymphocytes if appropriate immortalized human (or nonhuman) cell lines are available. For the purpose of the present invention, the animal of choice may be injected with an antigenic amount, for example, from about 0.1 mg to about 20 mg of the enzyme or protein or antigenic parts thereof. Usually the injecting material is emulsified in Freund's complete adjuvant. Boosting injections may also be required. The detection of antibody production can be carried out by testing the antisera with appropriately labelled antigen. Lymphocytes can be obtained by removing the spleen of lymph nodes of sensitized animals in a sterile fashion and carrying out fusion. Alternatively, lymphocytes can be stimulated or immunized in vitro, as described, for example, in Reading, Journal of Immunological Methods 53: 261-291, 1982.

A number of cell lines suitable for fusion have been developed and the choice of any particular line for hybridization protocols is directed by any one of a number of criteria such as speed, uniformity of growth characteristics, deficiency of its metabolism for a component of the growth medium, and potential for good fusion frequency.

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Intraspecies hybrids, particularly between like strains, work better than interspecies fusions. Several cell lines are available, including mutants selected for the loss of ability to secrete myeloma immunoglobulin.

Cell fusion can be induced either by virus, such as Epstein-Barr or Sendai virus, or polyethylene glycol. Polyethylene glycol (PEG) is the most efficacious agent for the fusion of mammalian somatic cells. PEG itself may be toxic for cells and various concentrations should be tested for effects on viability before attempting fusion. The molecular weight range of PEG may be varied from 1000 to 6000. It gives best results when diluted to from about 20% to about 70% (w/w) in saline or serum-free medium. Exposure to PEG at 37°C for about 30 seconds is preferred in the present case, utilizing murine cells. Extremes of temperature (i.e., about 45°C) are avoided, and preincubation of each component of the fusion system at 37°C prior to fusion can be useful. The ratio between lymphocytes and malignant cells is optimized to avoid cell fusion among spleen cells and a range of from about 1:1 to about 1:10 is commonly used.

The successfully fused cells can be separated from the myeloma line by any technique known by the art. The most common and preferred method is to choose a malignant line which is Hypoxanthine Guanine Phosphoribosyl Transferase (HGPRT) deficient, which will not grow in an aminopterin-containing medium used to allow only growth of hybrids and which is generally composed of hypoxanthine $1 \times 10^{-4} \text{M}$, aminopterin $1 \times 10^{-5} \text{M}$, and thymidine $3 \times 10^{-5} \text{M}$, commonly known as the HAT medium. The fusion mixture can be grown in the HAT-containing culture medium immediately after the fusion 24 hours later. The

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feeding schedules usually entail maintenance in HAT medium for two weeks and then feeding with either regular culture medium or hypoxanthine, thymidine-containing medium.

5 The growing colonies are then tested for the presence of antibodies that recognize the antigenic preparation. Detection of hybridoma antibodies can be performed using an assay where the antigen is bound to a solid support and allowed to react to
10 hybridoma supernatants containing putative antibodies. The presence of antibodies may be detected by "sandwich" techniques using a variety of indicators. Most of the common methods are sufficiently sensitive for use in the range of
15 antibody concentrations secreted during hybrid growth.

 Cloning of hybrids can be carried out after 21-23 days of cell growth in selected medium. cloning can be preformed by cell limiting dilution in fluid
20 phase or by directly selecting single cells growing in semi-solid agarose. For limiting dilution, cell suspensions are diluted serially to yield a statistical probability of having only one cell per well. For the agarose technique, hybrids are seeded
25 in a semi-solid upper layer, over a lower layer containing feeder cells. The colonies from the upper layer may be picked up and eventually transferred to wells.

 Antibody-secreting hybrids can be grown in
30 various tissue culture flasks, yielding supernatants with variable concentrations of antibodies. In order to obtain higher concentrations, hybrids may be transferred into animals to obtain inflammatory ascites. Antibody-containing ascites can be
35 harvested 8-12 days after intraperitoneal injection. The ascites contain a higher concentration of

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antibodies but include both monoclonals and immunoglobulins from the inflammatory ascites. Antibody purification may then be achieved by, for example, affinity chromatography.

5 The presence of the enzyme or protein contemplated herein in a species of Listeria may be accomplished in a number of ways such as on a semi-solid growth medium plate where colonies of bacteria are tested, by for example, the Western blotting
10 procedure. Alternatively, other semi-solid supports may be employed onto which the biological sample has been immobilized. In another method, the biological sample is contacted with a solid support already
15 containing a specific antibody. In any event the principle behind the assay is the same. A wide range of immunoassay techniques are available as can be seen by reference to US Patent No. 4,016,043, 4, 424,279 and 4,018,653. This, of course, includes
20 both single-site and two-site, or "sandwich", assays of the non-competitive types, as well as in the traditional competitive binding assays.

 Sandwich assays are among the most useful and commonly used assays and are favored for use in the present invention. A number of variations of the
25 sandwich assay technique exist, and all are intended to be encompassed by the present invention. Briefly, in a typical forward assay, an unlabelled antibody is immobilized on a solid substrate and the sample to be tested brought into contact with the bound molecule.
30 After a suitable period of incubation, for a period of time sufficient to allow formation of an antibody-antigen secondary complex, a second antibody specific to the antigen, labelled with a reporter molecule capable of producing a detectable signal is then
35 added and incubated, allowing time sufficient for the formation of a tertiary complex of antibody-antigen-

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labelled antibody. Any unreacted material is washed away, and the presence of the antigen is determined by observation of a signal produced by the reporter molecule. The results may either be qualitative, by
5 simple observation of the visible signal, or may be quantitated by comparing with a control sample containing known amounts of hapten. Variations on the forward assay include a simultaneous assay, in which both sample and labelled antibody are added
10 simultaneously to the bound antibody. These techniques are well known to those skilled in the art, including any minor variations as will be readily apparent.

In the typical forward sandwich assay, a first
15 antibody having specificity for the enzyme or protein, or antigenic parts thereof, contemplated in this invention, is either covalently or passively bound to a solid surface. The solid surface is typically glass or a polymer, the most commonly used
20 polymers being cellulose, polyacrylamide, nylon, polystyrene, polyvinyl chloride or polypropylene. The solid supports may be in the form of tubes, beads, discs of microplates, or any other surface suitable for conducting an immunoassay. The binding
25 processes are well-known in the art and generally consist of cross-linking covalently binding or physically adsorbing, the polymer-antibody complex is washed in preparation for the test sample. An aliquot of the sample to be tested is then added to
30 the solid phase complex and incubated at 25°C for a period of time sufficient to allow binding of any subunit present in the antibody. The incubation period will vary but will generally be in the range of about 2-40 minutes. Following the incubation
35 period, the antibody subunit solid phase is washed and dried and incubated with a second antibody

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specific for a portion of the hapten. The second antibody is linked to a reporter molecule which is used to indicate the binding of the second antibody to the hapten.

5 An alternative method involves immobilizing the target molecules in the biological sample and then exposing the immobilized target to specific antibody which may or may not be labelled with a reporter molecule. Depending on the amount of target and the
10 strength of the reporter molecule signal, a bound target may be detectable by direct labelling with the antibody. Alternatively, a second labelled antibody, specific to the first antibody is exposed to the target-first antibody complex to form a target-first
15 antibody-second antibody tertiary complex. The complex is detected by the signal emitted by the reporter molecule.

 By "reporter molecule" as used in the present specification, is meant a molecule which, by its
20 chemical nature, provides an analytically identifiable signal which allows the detection of antigen-bound antibody. Detection may be either qualitative or quantitative. The most commonly used reporter molecules in this type of assay are either
25 enzymes, fluorophores, luminescent molecules or radionuclide containing molecules (i.e. radioisotopes).

 In the case of an enzyme immunoassay, an enzyme is conjugated to the second antibody, generally by
30 means of glutaraldehyde or periodate. As will be readily recognized, however, a wide variety of different conjugation techniques exist, which are readily available to the skilled artisan. Commonly used enzymes include horseradish peroxidase, glucose
35 oxidase, beta-galactosidase and alkaline phosphatase, amongst others. The substrates to be used with the

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specific enzymes are generally chosen for the production, upon hydrolysis by the corresponding enzyme, of a detectable color change. For example, p-nitrophenyl phosphate is suitable for use with alkaline phosphatase conjugates; for peroxidase conjugates, 1,2-phenylenediamine, 5-aminosalicylic acid, or toluidine are commonly used. It is also possible to employ fluorogenic substrates, which yield a fluorescent product rather than the chromogenic substrates noted above. In all cases, the enzyme-labelled antibody is added to the first antibody hapten complex, allowed to bind, and then the excess reagent is washed away. A solution containing the appropriate substrate is then added to the tertiary complex of antibody-antigen-antibody. The substrate will react with the enzyme linked to the second antibody, giving a qualitative visual signal, which may be further quantitated, usually spectrophotometrically, to give an indication of the amount of hapten which was present in the sample. "Reporter molecule" also extends to use of cell agglutination or inhibition of agglutination such as red blood cells on latex beads, and the like.

Alternately, fluorescent compounds, such as fluorescein and rhodamine, may be chemically coupled to antibodies without altering their binding capacity. When activated by illumination with light of a particular wavelength, the fluorochrome-labelled antibody adsorbs the light energy, inducing a state to excitability in the molecule, followed by emission of the light at a characteristic color visually detectable with a light microscope. As in the EIA, the fluorescent labelled antibody is allowed to bind to the first antibody-hapten complex. After washing off the unbound reagent, the remaining tertiary complex is then exposed to the light of the

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appropriate wavelength the fluorescence observed indicates the presence of the hapten of interest. Immunofluorescent and EIA techniques are both very well established in the art and are particularly preferred for the present method. However, other reporter molecules, such as radioisotope, chemiluminescent or bioluminescent molecules, may also be employed. It will be readily apparent to the skilled technician how to vary the procedure to suit the required purpose.

Accordingly, an aspect of the present invention contemplates a method for detecting a species of the genus Listeria in a biological sample comprising the steps of contacting said biological sample with an antibody to an enzyme or other protein or parts thereof wherein said enzyme or other protein is involved in or associated with the metabolism of a carbon source in a species of Listeria and which is absent from another species of same genus for a time and under conditions sufficient for an enzyme- or other protein-antibody complex to form and subjecting said complex to a detecting means. The present invention contemplates that the antibody binding can be visualized by a number of methods including fluorescent staining.

In accordance with this method, the target bacteria or extracts thereof may first be immobilized onto a solid support. Where the target bacteria is immobilized, solid supports such as slides, plates, filters, beads or the like may be used. Where extracts such as protein are immobilized, solid supports such as resins can be used (resins are readily available commercially). On the other hand, protein may also be immobilized on supports such as those mentioned above for bacteria.

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The present invention is also directed to a kit for the rapid and convenient assay for species of Listeria in a biological sample. The kit is compartmentalized to receive a first container adapted to contain a biological sample to be tested and a second container adapted to contain an antibody to an enzyme or protein of species of Listeria as defined above, said antibody being labelled with a reporter molecule capable of giving a detectable signal as hereinbefore described. Alternatively, if said first antibody is not labelled then a further container is provided adapted to contain a second antibody to said first antibody where said second antibody is labelled with a reporter molecule. If the reporter molecule is an enzyme, then another container adapted to contain a substrate for said enzyme is provided.

Whether by a) genetic repair, b) hybridization, or c) antibody binding, the above three detection approaches require identification of enzymes, proteins or portions thereof, or identification of nucleic acid defining a promoter region, functional gene, regulatory gene or other control sequence or parts thereof for the expression of enzymes or other molecules involved in differential uptake or metabolism of the carbon sources of the present invention. Some enzymes involved in uptake or metabolism of some of these carbon sources are already known (e.g. L-lactic acid dehydrogenase, L-lactic acid oxidase, hexose phosphate phosphatase, etc.). Mutants deficient in some of these enzymes (and thus useful for transformation assays) are also known. See F. Mat-Jan et al., J. Bacterio. 171:342 (1989). Furthermore, nucleotide sequences of the genes for some enzymes (useful for the design of

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probes in hybridization assays) are described. E. Pradel et al., J. Bacterio. 172:802 (1990). Finally, such enzymes have been purified and antibodies have been raised against them (useful in immunological assays). M. Fatai and H. Kimura, J. Biol. Chem. 252:5820 (1977).

The present invention is further described by the following non-limiting Examples.

EXAMPLE ONE: Detection and Speciation by
Differential Carbon Source
Metabolism

Using the invention described herein, species of Listeria were screened for utilization of a variety of carbon sources on the Biolog MT MicroPlate™, a commercially available (Biolog, Inc., Hayward, California) 96-well microplate designed to test the ability of an inoculated microorganism suspension to utilize (oxidize) a panel of different carbon sources. Each well of the panel contains a tetrazolium redox dye and a buffered nutrient medium. A panel of carbon sources of the present invention was added to the MT plates and dried before inoculating with a microorganism suspension. Approximately 0.6 mg of each carbon source (e.g. 15 ul of a 4% stock solution) was added to each well. The Biolog MT MicroPlate™ was arranged so that there was one control well with no carbon source and the remaining wells contain individual carbon sources. The indicator plate was then incubated at an appropriate temperature of 37°C for 24 hours. Many of the carbon source reactions were apparent even by 1 hour. The results are shown in Table 1.

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TABLE 1
DIFFERENTIAL METABOLISM OF LISTERIA SPECIES AS
MEASURED BY REDOX DYE REDUCTION ON A
MICROPLATE™ TEST PANEL

SPECIES	D-GLUCONIC ACID L-MALIC ACID MANNITOL	5' AMP	GLYCEROL B-METHYL GLUCOSIDE	D-XYLOSE	L-RHAMNOSE	XYLITOL	SUCROSE	α-D- LACTOSE	L-LACTIC ACID	F6P G6P G1P	GENERALLY PATHOGENIC
MONOCYTOGENES	0	100	100	78 ^a	89 ^a	100	22	78 ^a	67	56	+
IVANOVII	0	100	100 ^a	100 ^a	0	0	0	100 ^a	100	100	+
INNOCUA	0	100	100	100 ^a	100 ^a	100	50 ^a	100 ^a	0	0	-
WELSHIMERI	0	100	100	100 ^a	50 ^a	100	50 ^a	100 ^a	0	0	-
SEELIGERI	0	100	100	100 ^a	0	100	0	100 ^a	0	0	-
GRAYI	100	0	100 ^a	0	0	0	0	0	0	0	-
MURRAYI	100	0	100	0	0	0	0	0	0	0	-

15 The numbers in the table indicate the percentage of strains of the particular species tested that could utilize the carbon source. Superscript "a" indicates reactions that are negative or very weak after 2 hours but positive at 24 hours. From the data it is

20 evident that some differences exist between the strains of a single species of Listeria.

The carbon sources L-lactic acid and hexose-phosphate (e.g. glucose-1-phosphate (G1P)) are of particular interest in this regard; these carbon

25 sources are utilized by some but not all of the strains currently regarded as monocytogenes and by most if not all strains of ivanovii. Since the overwhelming number of Listeria disease cases involve these species, the ability to distinguish strains of

30 monocytogenes and ivanovii offers the important potential to distinguish the pathogenic strains from the non-pathogenic strains.

In this regard, both L-lactic acid and hexose-phosphates are expected to be present in relatively

35 high levels inside the human body and if they can be utilized by the Listeria it would allow the Listeria to grow more easily inside the body. It has been recently shown that utilization of L-lactic acid by

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N. gonorrhoeae, in competition with leukocytes, gives it an advantage in escaping the normal defense processes of the human immune system. B.E. Britigan et al., J. Clin. Invest. 81:318 (1988). D.J. Hassett and M.S. Cohen, FASEB Journal 3:2574 (1989).

While there has heretofore been no suggestion that utilization of hexose phosphates is involved in pathogenicity, Listeria does grow as an intracellular pathogen and glucose (and other hexoses) are phosphorylated when transported into the cell. Therefore, the level of hexose phosphate should exceed the level of free hexose and provide a growth advantage to any invading bacteria capable of utilizing hexose phosphate as a carbon source.

Other links have been found between utilization of specific carbon source and pathogenicity of a certain species. This literature has been discussed in a recent review (Journal of General Microbiology, 1990 v. 136, p. 377). For example, Corynebacterium renale and Proteus mirabilis can grow in the kidneys and cause severe damage due, at least in part, to their synthesis of a urease enzyme which allows them to utilize urea. Another example is Brucella abortus which causes Brucellosis, resulting in fetal abortions in many animal species. Fetal tissues contain a high level of erythritol, and this bacterium is rather unusual in having the capability to utilize erythritol as a carbon source.

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EXAMPLE TWO: Strain Differentiation

Using the invention described herein, strains of species of Listeria were screened for growth on a variety of carbon sources on the Biolog MT MicroPlate™, a commercially available (Biolog, Inc., Hayward, California) 96-well microplate designed to test the ability of an inoculated microorganism suspension to utilize (oxidize) a panel of different carbon sources. As in Example 1, above, each well of the panel contains a tetrazolium redox dye and a buffered nutrient medium. A panel of carbon sources of the present invention was added to the MT plates and dried before inoculating with a microorganism suspension. About 0.6 mg of carbon source (e.g. 15 ul of a 4% stock solution) was added to each well. The Biolog's MT MicroPlate™ was arranged so that there was one control well with no carbon source and the remaining wells contain individual carbon sources. The indicator plate was then incubated at an appropriate temperature of 37°C for 24 hours. Some carbon source reactions were apparent at 1 hour. The results are shown in Table 2.

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TABLE 2
REACTIONS OF STRAINS OF LISTERIA SPECIES IN
BIOLOG'S TEST PANEL

5	SPECIES ^b	STRAIN #	D-GLUCONIC ACID L-MALIC ACID		GLYCEROL B-METHYL		D-XYLOSE	XYLITOL	L-LACTIC ACID		F6P G6P		α-D-
			MANNITOL	5' AME	GLUCOSIDE	L-RHAMNOSE					G1P	SUCROSE	
10	MONOCYTOGENES	1293	-	+	+	+ ^a	-	+	-	-	+	+	+
		2128	-	+	+	-	+ ^a	+	+	+	-	-	+ ^a
		2144	-	+	+	+ ^a	+ ^a	+	+	+ ^a	+	-	-
		2198	-	+	+	+ ^a	+ ^a	+	-	-	-	-	+ ^a
15	IVANOVII	8276	-	+	+	+ ^a	+ ^a	+	-	-	-	-	+ ^a
		8277	-	+	+	+ ^a	+ ^a	+	+	+	-	-	+ ^a
		8278	-	+	+	+ ^a	+ ^a	+	+	+ ^a	-	-	+ ^a
		8279	-	+	+	+ ^a	+ ^a	+	+	+	-	-	+ ^a
20	WELSHIMERI	8280	-	+	+	+ ^a	-	+	+ ^a	-	-	-	-
		2103	-	+	+ ^a	-	+ ^a	-	+	+	-	-	+ ^a
		9076	-	+	+ ^a	-	+ ^a	-	+	+	-	-	+ ^a
		2102	-	+	+	+ ^a	+ ^a	+	-	-	-	-	+ ^a
25	GRAYI	9075	-	+	+	+ ^a	+ ^a	+	-	-	+ ^a	-	+ ^a
		2105	-	+	+	+ ^a	+ ^a	+	-	-	-	-	+ ^a
		9079	-	+	+	-	+ ^a	+	-	-	+ ^a	-	+ ^a
		2104	-	+	+	-	+ ^a	+	-	-	-	-	+ ^a
30	MURRAYI	9078	-	+	+	-	+ ^a	+	-	-	-	-	-
		2101	+	-	+ ^a	-	-	-	-	-	-	-	-
		9074	+	-	+ ^a	-	-	-	-	-	-	-	-
		2199	+	-	+	-	-	-	-	-	-	-	-
		9077	+	-	+	-	-	-	-	-	-	-	-

^a Negative or very weak reaction after 2 hours.^b These are the species designations on the strain when received.

30 It is clear from Table 2 that the method of the present invention is able to differentiate between strains of particular Listeria species. The source of the strains is provided in Table 3.

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TABLE 3

CHARACTERIZATION OF STRAINS OF SPECIES OF LISTERIA

	STRAIN #	SPECIES	SOURCE ¹	SOURCE #
5	2101	GRAYI	CCUG	EF-4983
	9074	GRAYI	ATCC	19120
	2102	INNOCUA	FDA	L-0705
	9075	INNOCUA	ATCC	33090
	2103	IVANOVII	FDA	KC-1714
10	9076	IVANOVII	ATCC	19119
	1293	MONOCYTOGENES	Carey	
	2128	MONOCYTOGENES	ATCC	15313
	2144	MONOCYTOGENES	U of IL	
	2198	MONOCYTOGENES	CCUG	1452
15	8276	MONOCYTOGENES	Iowa St.	VM-4
	8277	MONOCYTOGENES	CCUG	EF-440
	8278	MONOCYTOGENES	CCUG	EF-441
	8279	MONOCYTOGENES	CCUG	EF-442
	8280	MONOCYTOGENES	CCUG	EF-444
20	2199	MURRAYI	CCUG	EF-4984
	9077	MURRAYI	ATCC	25401
	2104	SEELIGERI	FDA	CA-0705
	9078	SEELIGERI	ATCC	35967
	2105	WELSHIMERI	FDA	CU-0705
	9079	WELSHIMERI	ATCC	35897
25	1 ATCC	=	American Type Culture Collection (type strains)	
	CCUG	=	Culture Collection University of Gothenburg, Sweden, Enevold Falsen	
	FDA	=	Food & Drug Administration Lab, Minneapolis MN, Sallie McLaughlin	
	IOWA ST.	=	Iowa State University, Paul Hartman	
	CAREY	=	Roberta Carey, St. Francis Hospital, Evanston, IL	
30	U of IL	=	University of Illinois, Chicago Med Center, Paula Malloy	

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WHAT IS CLAIMED IS :

1. A method for detecting a species of the genus Listeria comprising:

- 5 a) exposing a sample suspected of containing microorganisms belonging to the genus Listeria to a metabolically effective amount of one or more carbon sources capable of being metabolized by a subset of species of Listeria, for a time and under conditions sufficient for said subset
10 to metabolize said carbon source; and
b) detecting any metabolism of said carbon source by assaying for respiration by said microorganisms, thereby determining the presence of said subset of species.
- 15 2. The method of claim 1 further comprising, prior to said exposing step, subjecting said sample to an enrichment treatment which will enhance the population of the genus Listeria in said sample.
- 20 3. The method of claim 1 wherein said one or more carbon sources are selected from the group consisting of mannitol, D-gluconic acid, L-malic acid, 5'AMP, β -methyl glucoside, glycerol, D-xylose, L-rhamnose, xylitol, sucrose, α -D-lactose, L-lactic acid, and hexose-phosphate.
- 25 4. The method of claim 1 wherein said one or more carbon sources comprise a panel of carbon sources comprised of L-lactic acid and xylitol.
- 30 5. The method of claim 1 wherein said one or more carbon sources comprise a panel of carbon sources comprised of hexose-phosphate and xylitol.

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6. A test kit for the detection of a species of the genus Listeria or a strain of a species of the genus Listeria comprising at least one compartment containing at least one carbon source capable of being metabolized by a subset of species of the genus Listeria, wherein said carbon source is selected from the group consisting of mannitol, D-gluconic acid, L-malic acid, 5'AMP, β -methyl glucoside, glycerol, D-xylose, L-rhamnose, xylitol, sucrose, α -D-lactose, L-lactic acid, and hexose-phosphate.

7. A test kit for the detection of a species of the genus Listeria or a strain of a species of the genus Listeria comprising a plurality of compartments containing a plurality of carbon sources capable of being metabolized by a subset of species of the genus Listeria, wherein said carbon sources comprise a panel of carbon sources, each carbon source of said panel in separate compartments, comprising L-lactic acid and xylitol.

8. A test kit for the detection of a species of the genus Listeria or a strain of a species of the genus Listeria comprising a plurality of compartments containing a plurality of carbon sources capable of being metabolized by a subset of species of the genus Listeria, wherein said carbon sources comprise a panel of carbon sources, each carbon source of said panel in separate compartments, comprising hexose-phosphate and xylitol.

9. A method for detecting a species of the genus Listeria comprising:
a) exposing a sample suspected of containing microorganisms belonging to the genus Listeria to a metabolically effective amount of one or

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5 more carbon sources capable of being metabolized by L. monocytogenes and L. ivanovii but not for metabolism by other species of Listeria, for a time and under conditions sufficient for metabolism by L. monocytogenes and L. ivanovii ; and

b) detecting any metabolism of said carbon source, thereby determining the presence of L. monocytogenes and/or L. ivanovii.

10 10. The method of claim 9 further comprising, prior to said exposing step, subjecting said sample to an enrichment treatment which will enhance the population of the genus Listeria in said sample.

15 11. The method of claim 9 wherein said one or more carbon sources are selected from the group consisting of L-lactic acid and hexose-phosphate.

12. A method for detecting a species of the genus Listeria comprising:

20 a) exposing a sample suspected of containing microorganisms belonging to the genus Listeria to a metabolically effective amount of one or more carbon sources capable of being metabolized by a subset of species of the genus Listeria, but not metabolized by L. monocytogenes or L. ivanovii, for a time and under conditions sufficient for metabolism by said subset of species; and

25 b) detecting any metabolism of said carbon source, thereby determining the presence of a species other than L. monocytogenes and L. ivanovii.
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13. The method of claim 12 further comprising, prior to said exposing step, subjecting said sample to an enrichment treatment which will enhance the population of the genus Listeria in said sample.
- 5 14. The method of claim 12 wherein said one or more carbon sources are selected from the group consisting of mannitol, D-gluconic acid, and L-malic acid.
- 10 15. The method of claim 12 wherein said one or more carbon sources comprise a panel of carbon sources comprised of mannitol, D-gluconic acid, and L-malic acid.
- 15 16. An enrichment medium for species of the genus Listeria comprising, in liquid or semi-solid form:
a basal medium having all the constituents selective for growth of the genus Listeria;
and
a growth effective amount of one or more carbon sources capable of being utilized by a subset of species of the genus Listeria, wherein the
20 carbon source is selected from the group consisting of mannitol, D-gluconic acid, L-malic acid, 5'AMP, β -methyl glucoside, glycerol, D-xylose, L-rhamnose, xylitol, sucrose, α -D-lactose, L-lactic acid, and hexose-phosphate.
- 25 17. In a method for detecting microorganisms of the genus Listeria wherein a sample suspected of containing microorganisms belonging to the genus Listeria is subjected to an enrichment treatment capable of enhancing the population of the genus
30 Listeria in said sample, the improvement comprising:
a) thereafter exposing said sample to a metabolically effective amount of one or more

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carbon sources capable of being metabolized by a subset of species of Listeria, for a time and under conditions sufficient for said subset to metabolize said carbon source; and

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b) detecting any metabolism of said carbon source by assaying for respiration by said microorganisms, thereby determining the presence of said subset of species.

INTERNATIONAL SEARCH REPORT

International Application No. PCT/US91/04699

I. CLASSIFICATION OF SUBJECT MATTER (if several classification symbols apply, indicate all) ⁶		
According to International Patent Classification (IPC) or to both National Classification and IPC		
IPC(5): C12Q 1/02		
U.S. CL.: 435/29		
II. FIELDS SEARCHED		
Minimum Documentation Searched ⁷		
Classification System	Classification Symbols	
U.S.	435/29, 822; 424/7.1; 436/63	
Documentation Searched other than Minimum Documentation to the extent that such Documents are included in the Fields Searched ⁸		
BIOSIS: "AUXOTRO?" "LISTERIA"		
III. DOCUMENTS CONSIDERED TO BE RELEVANT ⁹		
Category [*]	Citation of Document, ¹¹ with indication, where appropriate, of the relevant passages ¹²	Relevant to Claim No. ¹³
A	Journal of General Microbiology, Volume 98, issued 1977, B.A. WILKINSON ET AL., "A Numerical Taxonomic Survey of Listeria and Related Bacteria", pages 399-421, see entire document.	1-15, 17
Y	SNEATH ET AL., "Bergey's Manual of Systematic Bacteriology, Volume 2," published 1986 by Williams & Wilkins, see pages 1235-1249.	1-15, 17
Y	Applied and Environmental Microbiology, Volume 33, No. 2, issued February 1977, B.R. BOCHNER ET AL., "Generalized Indicator Plate for Genetic, Metabolic, and Taxonomic Studies with Microorganisms", pages 434-444, see entire document.	1-15, 17
Y	Biological Abstracts, issued 1977, VISWANATH ET AL., "Characterization of Glycerol Nonutilizing and Proto-perithecial Mutants of Neurospora, see Abstract No. 5021.	3-16
(cont.)		
<p>[*] Special categories of cited documents: ¹⁰</p> <p>"A" document defining the general state of the art which is not considered to be of particular relevance</p> <p>"E" earlier document but published on or after the international filing date</p> <p>"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)</p> <p>"O" document referring to an oral disclosure, use, exhibition or other means</p> <p>"P" document published prior to the international filing date but later than the priority date claimed</p> <p>"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention</p> <p>"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step</p> <p>"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.</p> <p>"&" document member of the same patent family</p>		
IV. CERTIFICATION		
Date of the Actual Completion of the International Search		Date of Mailing of this International Search Report
19 September 1991		25 OCT 1991
International Searching Authority		Signature of Authorized Officer
ISA/US		Jane Williams (vsh)

Form PCT/ISA/210 (second sheet) (Rev.11-87)

FURTHER INFORMATION CONTINUED FROM THE SECOND SHEET

A	Biological Abstracts, issued 15 May 1976, SPRAGE ET AL. , "A Mutant of Escherichia coli auxotrophic for organic phosphates: Evidence for 2 defects in inorganic phosphate transport," see Abstract No. 55005.	3,5,6,8,11,16
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V. ☐ OBSERVATIONS WHERE CERTAIN CLAIMS WERE FOUND UNSEARCHABLE¹

This international search report has not been established in respect of certain claims under Article 17(2) (a) for the following reasons:

1. ☐ Claim numbers because they relate to subject matter ¹² not required to be searched by this Authority, namely:

2. ☐ Claim numbers because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out ¹³, specifically:

3. ☐ Claim numbers because they are dependent claims not drafted in accordance with the second and third sentences of PCT Rule 6.4(a).

VI. ☐ OBSERVATIONS WHERE UNITY OF INVENTION IS LACKING²

This International Searching Authority found multiple inventions in this international application as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims of the international application.
2. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims of the international application for which fees were paid, specifically claims:
3. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claim numbers:
4. ☐ As all searchable claims could be searched without effort justifying an additional fee, the International Searching Authority did not invite payment of any additional fee.

Remark on Protest

- ☐ The additional search fees were accompanied by applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.